| Bparg | 1 | MSAGPGCEPCTKRPRWDAAATSPPAASDARSFPGRQRRVLDSKDAPVQFRVPPSSSGCALGRAGQERGSATSL |
| Eparg | 1 | MNAGPGCEPCTKATRWGAATTS.PAASDARSFPSRQRRVLDPKDAEVQFRVPPSSPACVPGQAGQERGSATSL |
| Mparg | 1 | MSAGPGWEPCTKA.RWGAAGTSAPIASDARSFPGRQRRVLDPKDAPVQFRVPPSSPACVSGRAGPERGSATHF |
| Dparg | 1 | ------------- |
| Ceparg1 | | ------------- |

| Bparg | 74 | VFKQKTITSWMDTKGIKTVESKSLESXEHHHTREESMMSSVQKDHFYQHHMEKLEHVSQLGFDKSPVEKGTQY |
| Eparg | 73 | VFKQKTITSWMDTKGIKTVESKSLDSXEHHHTREESMMSSVQKDHFYQHHVEKLVHVSQLSLDKSLTEKSTQY |
| Mparg | 73 | VFKQKTITSWMDTKGIKTVESK...SXEHHHTREESMMSSVQKDHFYPHKVEKLEHVPQLNLDKSPTEKSSQY |
| Dparg | 1 | ------------- |
| Ceparg1 | | ------------- |

| Bparg | 218 | LKQHQTAAMCKWQHHGPHSERLLESEPPAVTLVPEQFSHAHVDQSSPKDDHSDTNSEESRDNQQFLTHVKL.. |
| Eparg | 217 | LKQHQTAAMCKWQHHGKHTEQLLESEPQTVTLVPEQFSHAHIDRSSPQNDHSDTDSEEHRDNQQFLTVKL.. |
| Mparg | 214 | LKQQQTASVCKWQHHGKHAEQLLASEPPAGTPLPKQLSHAHIGQSPETDDHSDTDHEEDRDNQQFLTPIKL.. |
| Dparg | 1 | ------------- |
| Ceparg1 | | MSKKFIELGDPVTQDEKDYEDYVGVGFAEQVP |

| Bparg | 298 | .ANAKQTMEDEQGREARSHQKCGKACHPAEACAGCQQEETDVVSESPLSDTGSEDVGTGLKHAHRLHRQESSL |
| Eparg | 297 | .ANAKQTEDEHAREAKSHQKCSKSCHPGEDCASCQQDEIDVVPKSPLSDVGSEDVGTGSKHDHKLIRQESCL |
| Mparg | 291 | .ANTKPTVGDGQ...ARSHCKCSGSRQSKEDCTGCQQEEVDVLPESPLSDVGAEDIGTGPKHDHKLTGQESSL |
| Dparg | 1 | ------------- |
| Ceparg | 33 | TMKRRKLTEEGNTTESKEDPEEPKSRDVFVSSQSSDESQEDSAENPEIAKEVSENCENLTETLKISNIESLDN |

| Bparg | 396 | GHSPPFEKESEPESPMDVDNSKHSCQDSEADEETSPGFDE.QEDSSSAQTANKPSRFQPREADTELRKRSSAK |
| Eparg | 395 | GHSPPFEKESEPESPMDVDNSKHSCQDSEADEETSPGFDE.QEDGSSSQTANKPSRFQARDADIEFRKRYSTK |
| Mparg | 387 | GDSPPFEKESEPESPMDVDNSKHSCQDSEADEETSPvFDE.QEDRSS.QTANKLSSCQAREADGDLRKRYLTK |
| Dparg | 1 | MQEFRSHLIFPIFQ.KVIQS.TANRRR.ASASVL.....TNRLGK..ALCLNCARMSKSPDGGISEIE... |
| Ceparg106 | | VTERSEHTLDN...HKSTEPHEE.DVNNKSHIDVAINSDEDDELVLEENHKEMRDGEQVQQLS..QDLFADD |

FIG. 16A

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Bparg   468  GGEIRLHFQFEGGESRAGMN.DVNAKRPGSTSSLNVECRNSKQEGRKDSKITDEFMRVPKAEDKRKEQCEMKH
Eparg   467  GGEVRLHFQFEGGESRTGMN.DLNAKLPGNISSLNVECRNSKQEGKKDSKITDELMRLPKAEDRRKEQWETKH
Mparg   354  GGEVRLHFQFE.GEMNAGTS.DLNAKPSGNSSSLNVECRSSKQEGKRDSKITDEFMRISKSEDRRKEQCEVRH
Dparg   59   .TEEE.PENLANSL..DDSWRGVSMEAIHRNRQPFELENLPPVTAGNLHRVMIQLPIRET..PPR.PIKSPG
Ceparg  172  QELIEIPGIMKDTTTQLDITDSEVETAQKMEMIEETEADSTFVGEDSKATKTVRTSSSSF...LSTVSTCEAP

Bparg   540  QRTERKIPKYYPPHLSPDKKWLGTPIEEM..RRMPRCGIRLPPLRPSANHTVTIRVDLLRIGEVPK.PFPTHF
Eparg   539  QRTERKIPKYVPPELSPDKKWLGTPIEEM..RRMPRCGIRLPLLRPSANHTVTIRVDLLRAGEVPK.PFPTHY
Mparg   425  QRTERKIPKYIPPMLPPEKKWLGTPIEEM..RKMPRCGIHLPSLRPSASHTVTVRVDLLRAGEVPK.PFPTHY
Dparg   125  X..WDSEEVRLPCAPESKYPRENPDGS.TTIDFRWEMIERALLQPIKTCEELQAAIISIHTTIRDQWEFRALE
Ceparg  242  AKGRARMIQKELEKEVIAFTEGNLT.....LQPDLNKVDPDRNIRI.CTIPNFPASQG..KLRED.NRIG..

Bparg   610  KDLWDNKEVXMPCSEQNLYPVEDENGE.RAAGSRWELIQTALLHRLTRPQNLKDAILKINVAISKKWDFTALI
Eparg   609  KDLWDNKEVXMPCSEQNLYPVEDENGE.RTAGSRWELIQTALLHKFTRPQNLKDAILKINVAISKKWDFTALI
Mparg   495  KDLWDNKEVXMPCSEQNLYPVEDENGE.RTAGSRWELIQTALLHKFTRPQNLKDAILKINVAISKKWDFTALV
Dparg   196  QLLDEELDESETRVFFQSILPDMVKIALCLPNICTQPIPLLKQKMNHSITMSQEQIASLLANAFFCTFPRRNA
Ceparg  367  PKIVLPQRWRE.FDSRGRR..RDSYFYFKRKLDGILKCIKTTGYFMFVGLLH......HMWE..FDPDITIK

Bparg   682  DFWDXVLEEAEAQHLYQSILPDMVKIALCLPNICTQPIPLLKQKMNHSITMSQEQIASLLANAFFCTFPRRNA
Eparg   681  DFWDXVLEEAEAQHLYQSILPDMVKIALCLPNICTQPIPLLKQKMNHSITMSQEQIASLLANAFFCTFPRRNA
Mparg   567  DFWDXVLEEAEAQHLYQSILPDMVKIALCLPNICTQPIPLLKQKMNHSVTMSQEQIASLLANAFFCTFPRRNA
Dparg   269  .KMKSEISSYPDIHFNRLFEGRSSRKPEKLKTLFCIFRRV.TERDASNVPTGVVTFVRRSGLPEE.L.IDWS
Ceparg  424  ....LPAL.EMYYKEMSELVGREEVLEKFARVARIAKTAEDILPERIYRLVGDVE.SATLSHKQCAALVARMF

Bparg   755  .KMKSEISSYPDIHFNRLFEGRSSRKPEKLKTLFCIFRRV..TEKK....PTGLVTFIRQS.L.ED.F.PEWE
Eparg   754  .KMKSEISSYPDIHFNRLFEGRSSRKPEKLKTLFCIFRRV..TEKK...PTGLVTFIRQS.L.ED.F.PEWE
Mparg   387  .KMKSEISSYPDIHFNRLFEGRSSRKPEKLKTLFCIFRRV..TEKK...PTGLVTFIRQS.L.ED.F.PEWE
Dparg   337  QSAAPLGDVPLEVDAEGTIEDEGIGLLQVDFAHKYLGGGVLGHGCVQEEIRFVICPELLVGKLFTECLRPFEA
Ceparg  491  FA.......RPDSPFS......FCRILSSDKSICVEKLKFLFTIF.....DKMSMDPPDGAVSF.RLTKMD
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# FIG. 16B

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Bparg   817  RCEKLL..TRLEVIYEGTIEGNGQGMLQVDFAHRFVGGGVTSAGLVQEEIRFLIHPELIVSRLFIEVLDHNEC
Eparg   816  RCEKPL..TRLEVIYEGTIEENGQGMLQVDFAHRFVGGGVTSAGLVQEEIRFLIHPELIISRLFIEVLDHNEC
Mparg   702  RCEKPL..TRLEVIYEGTIEGNGRGMLQVDFAHRFVGGGVTGAGLVQEEIRFLIHPELIVSRLFIEVLDHNEC
Dparg   410  LVMLGAERYSNYTGIAGSFEWS.....GNFEDSTP.RDSSGRRQTAIVAIDALHFA.QSHH....QYREDL
Ceparg  543  KDIFNEEW.XDXXLRSLPEVEFFDEMLIEDTAL.CTQVDFAHEHLGGGVLNHGSVQEEIRFLMCPEMMVGMLL

Bparg   888  LIITGTEQISEYTGIAETYRWA......RSHEDRSE.RDDWQRRTEIVAIDALHFR.RYLD....QFVPEKIR
Eparg   887  LIITGTEQISEYTGIAETYRWS......RSHEDRSE.RDDCERRCTEIVAIDALHFR.RYLD....QFVPEKMR
Mparg   773  LIITGTEQISEYTGIAETYRWA......RSHEDRSE.KDDWQRRCTEIVAIDALHFR.RYLD....QFVPEKVR
Dparg   472  MERELHKAYIGFVHWMVTPP..PGVATGNWGCGAFGGDSYLKALLQLMVCAQLGRPLAYYTFGNVEFRDDF
Ceparg  614  CEKMKQLEAISIVGAYVFSSYTGYGHTLKWAELQPNHSRQNTNEFRDRFGRLRVETIAIDAILFKGSKLDCQT

Bparg   951  RELHKAYCGFLRPGVSSEHLSAVATGNWGCGAFGGDARLKALIQYLAAAVAERDVVIFTFGDSELMRDIYSMH
Eparg   950  RELHKAYCGFLRPGVSSEHLSAVATGNWGCGAFGGDARLKALIQYLAAAVAERDVVIFTFGDSELMRDIYSMH
Mparg   835  RELHKAYCGFLRPGVPSEHLSAVATGNWGCGAFGGDARLKALIQYLAAAAAERDVVIFTFGDSELMRDIYSMH
Dparg   543  HEMWLLFRNDGTTVQQ.LWS.ILRSYSRLIKEKSSKEPRENKASKKKLYDFI...KEELKKVRDVPGEGAS
Ceparg  687  EQLNKANIIREMKKASIGFMSQGPKFTHIP.IVTGWWGCGAFNGDKPLKFIIQVIAAGVADRPLHFCSFGEPE

Bparg  1021  TFLTERKLTVGE.VYKLLRYNEECRNCSTPGP......DIKLYPFIYHAVESCTQTTNQPGQRTGA-------
Eparg  1020  IFLTERKLTVGD.VYKLLLRYNEECRNCSTPGP......DIKLYPFIYHAVESCAETADHSGQRTGT------
Mparg   907  TFLTERKLDVGK.VYKLLLRYNEECRNCSTPGP......DIKLYPFIYHAVESSAETTDMPGQKAGT------
Dparg   611  AEAGSSRVAGLGEGKSETSAKSSPELNKQPARPQITITQQSTDLLPAQLSQDNSNSSEDQALLMLSDDEEA
Ceparg  759  LAAXCKKIIERMKQKDVTLGKSCFSIFS-----

Bparg  1087  -------
Eparg  1086  -------
Mparg   969  -------
Dparg   684  NAMMEAASLEAKSSVEISNSSTTSKTSSTATKSMGSGGRQLSLLEMLDTHYEKGSASKRPRKSPNCSKAEG
Ceparg  787  -------
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# FIG. 16C

FIG. 16D